

SEQUENCE LISTING

<110> Coverley, Dawn

<120> REPLICATION PROTEIN

<130> 9052-222

<140> PCT/GB03/05334

<141> 2003-12-05

<150> GB 0228337.2

<151> 2002-12-05

<160> 73

<170> PatentIn version 3.1

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<212> PRT

<213> Homo sapiens

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Asp Ser Ser Ser Gln  
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<213> Homo sapiens

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Val Glu Glu Glu Leu Cys Lys Gln  
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ccggcttggt acagatat

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32

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tcttgcgatt gggggcggcg gaagagggg tt

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aagcagacac agggcccgga tcggctgcct

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aatcgcaagg attcttcttc tcctgtctc

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aaagaagaag aatccttgcg acctgtctc

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aatctgcagc agttctttcc ccctgtctc

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tccgagccct tccactcctc tctgg 25

<210> 26  
<211> 845  
<212> PRT  
<213> Mouse

<400> 26

Met Phe Asn Pro Gln Leu Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln  
1 5 10 15

Gln Gln Leu Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln  
20 25 30

Gln Gln Ile Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln  
35 40 45

Ala Ser Leu Ser Ile Pro Val Ser Arg Gly Leu Pro Gln Gln Ser Ser  
50 55 60

Pro Gln Gln Leu Leu Ser Leu Gln Gly Leu His Ser Thr Ser Leu Leu  
65 70 75 80

Asn Gly Pro Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly  
85 90 95

Leu Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Gly Ala Ser Leu  
 100 105 110

Thr Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Ala Phe Asn Val Thr  
 115 120 125

Ala Pro Ser Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Met Val Thr  
 130 135 140

Pro Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu  
 145 150 155 160

Gly Pro Pro Pro Val Gly Val Pro Ile Asn Pro Ser Gln Leu Asn His  
 165 170 175

Ser Gly Arg Asn Thr Gln Lys Gln Ala Arg Thr Pro Ser Ser Thr Thr  
 180 185 190

Pro Asn Arg Lys Asp Ser Ser Ser Gln Thr Val Pro Leu Glu Asp Arg  
 195 200 205

Glu Asp Pro Thr Glu Gly Ser Glu Glu Ala Thr Glu Leu Gln Met Asp  
 210 215 220

Thr Cys Glu Asp Gln Asp Ser Leu Val Gly Pro Asp Ser Met Leu Ser  
 225 230 235 240

Glu Pro Gln Val Pro Glu Pro Glu Pro Phe Glu Thr Leu Glu Pro Pro  
 245 250 255

Ala Lys Arg Cys Arg Ser Ser Glu Glu Ser Thr Glu Lys Gly Pro Thr  
 260 265 270

Gly Gln Pro Gln Ala Arg Val Gln Pro Gln Thr Gln Met Thr Ala Pro  
 275 280 285

Lys Gln Thr Gln Thr Pro Asp Arg Leu Pro Glu Pro Pro Glu Val Gln  
 290 295 300

Met Leu Pro Arg Ile Gln Pro Gln Ala Leu Gln Ile Gln Thr Gln Pro  
 305 310 315 320

Lys Leu Leu Arg Gln Ala Gln Thr Gln Thr Ser Pro Glu His Leu Ala  
 325 330 335  
 Pro Gln Gln Asp Gln Val Glu Pro Gln Val Pro Ser Gln Pro Pro Trp  
 340 345 350  
 Gln Leu Gln Pro Arg Glu Thr Asp Pro Pro Asn Gln Ala Gln Ala Gln  
 355 360 365  
 Thr Gln Pro Gln Pro Leu Trp Gln Ala Gln Ser Gln Lys Gln Ala Gln  
 370 375 380  
 Thr Gln Ala His Pro Gln Val Pro Thr Gln Ala Gln Ser Gln Glu Gln  
 385 390 395 400  
 Thr Ser Glu Lys Thr Gln Asp Gln Pro Gln Thr Trp Pro Gln Gly Ser  
 405 410 415  
 Val Pro Pro Pro Glu Gln Ala Ser Gly Pro Ala Cys Ala Thr Glu Pro  
 420 425 430  
 Gln Leu Ser Ser His Ala Ala Glu Ala Gly Ser Asp Pro Asp Lys Ala  
 435 440 445  
 Leu Pro Glu Pro Val Ser Ala Gln Ser Ser Glu Asp Arg Ser Arg Glu  
 450 455 460  
 Ala Ser Ala Gly Gly Leu Asp Leu Gly Glu Cys Glu Lys Arg Ala Gly  
 465 470 475 480  
 Glu Met Leu Gly Met Trp Gly Ala Gly Ser Ser Leu Lys Val Thr Ile  
 485 490 495  
 Leu Gln Ser Ser Asn Ser Arg Ala Phe Asn Thr Thr Pro Leu Thr Ser  
 500 505 510  
 Gly Pro Arg Pro Gly Asp Ser Thr Ser Ala Thr Pro Ala Ile Ala Ser  
 515 520 525  
 Thr Pro Ser Lys Gln Ser Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala  
 530 535 540  
 Ser Ser Ser Ser Gln Gln Glu Phe Gln Asp His Met Ser Glu Ala Gln



545					550					555					560
His	Gln	Gln	Arg	Leu	Gly	Glu	Ile	Gln	His	Ser	Ser	Gln	Thr	Cys	Leu
				565					570					575	
Leu	Ser	Leu	Leu	Pro	Met	Pro	Arg	Asp	Ile	Leu	Glu	Lys	Glu	Ala	Glu
			580					585					590		
Asp	Pro	Pro	Pro	Lys	Arg	Trp	Cys	Asn	Thr	Cys	Gln	Val	Tyr	Tyr	Val
		595					600					605			
Gly	Asp	Leu	Ile	Gln	His	Arg	Arg	Thr	Gln	Glu	His	Lys	Val	Ala	Lys
610						615					620				
Gln	Ser	Leu	Arg	Pro	Phe	Cys	Thr	Ile	Cys	Asn	Arg	Tyr	Phe	Lys	Thr
625					630					635					640
Pro	Arg	Lys	Phe	Val	Glu	His	Val	Lys	Ser	Gln	Gly	His	Lys	Asp	Lys
				645					650					655	
Ala	Gln	Glu	Leu	Lys	Thr	Leu	Glu	Lys	Glu	Thr	Gly	Ser	Pro	Asp	Glu
			660					665						670	
Asp	His	Phe	Ile	Thr	Val	Asp	Ala	Val	Gly	Cys	Phe	Glu	Ser	Gly	Gln
		675					680					685			
Glu	Glu	Asp	Glu	Asp	Asp	Asp	Glu	Glu	Glu	Glu	Glu	Glu	Gly	Glu	Ile
690						695						700			
Glu	Ala	Glu	Glu	Glu	Phe	Cys	Lys	Gln	Val	Lys	Pro	Arg	Glu	Thr	Ser
705					710					715					720
Ser	Glu	Gln	Gly	Lys	Gly	Ser	Glu	Thr	Tyr	Asn	Pro	Asn	Thr	Ala	Tyr
				725					730					735	
Gly	Glu	Asp	Phe	Leu	Val	Pro	Val	Met	Gly	Tyr	Val	Cys	Gln	Ile	Cys
			740					745					750		
His	Lys	Phe	Tyr	Asp	Ser	Asn	Ser	Glu	Leu	Arg	Leu	Ser	His	Cys	Lys
		755					760					765			
Ser	Leu	Ala	His	Phe	Glu	Asn	Leu	Gln	Lys	Tyr	Lys	Ala	Lys	Asn	Pro
770						775						780			

Ser Pro Pro Pro Thr Arg Pro Val Ser Arg Lys Cys Ala Ile Asn Ala  
785 790 795 800

Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser Ser His Gln Pro Ser Pro  
805 810 815

Gln Asp Thr Val Lys Met Pro Ser Lys Val Lys Pro Gly Ser Pro Gly  
820 825 830

Leu Pro Pro Pro Leu Arg Arg Ser Thr Arg Leu Lys Thr  
835 840 845

<210> 27  
<211> 716  
<212> PRT  
<213> Mouse

<400> 27

Ser Thr Ser Leu Leu Asn Gly Pro Met Leu Gln Arg Ala Leu Leu Leu  
1 5 10 15

Gln Gln Leu Gln Gly Leu Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr  
20 25 30

Asp Gly Ala Ser Leu Thr Met Pro Thr Ala Thr Leu Gly Asn Leu Arg  
35 40 45

Ala Phe Asn Val Thr Ala Pro Ser Leu Ala Ala Pro Ser Leu Thr Pro  
50 55 60

Pro Gln Met Val Thr Pro Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr  
65 70 75 80

Arg Gln Ser Leu Leu Gly Pro Pro Pro Val Gly Val Pro Ile Asn Pro  
85 90 95

Ser Gln Leu Asn His Ser Gly Arg Asn Thr Gln Lys Gln Ala Arg Thr  
100 105 110

Pro Ser Ser Thr Thr Pro Asn Arg Lys Thr Val Pro Leu Glu Asp Arg  
115 120 125

Glu Asp Pro Thr Glu Gly Ser Glu Glu Ala Thr Glu Leu Gln Met Asp  
 130 135 140

Thr Cys Glu Asp Gln Asp Ser Leu Val Gly Pro Asp Ser Met Leu Ser  
 145 150 155 160

Glu Pro Gln Val Pro Glu Pro Glu Pro Phe Glu Thr Leu Glu Pro Pro  
 165 170 175

Ala Lys Arg Cys Arg Ser Ser Glu Glu Ser Thr Glu Lys Gly Pro Thr  
 180 185 190

Gly Gln Pro Gln Ala Arg Val Gln Pro Gln Thr Gln Met Thr Ala Pro  
 195 200 205

Lys Gln Thr Gln Thr Pro Asp Arg Leu Pro Glu Pro Pro Glu Val Gln  
 210 215 220

Met Leu Pro Arg Ile Gln Pro Gln Ala Leu Gln Ile Gln Thr Gln Pro  
 225 230 235 240

Lys Leu Leu Arg Gln Ala Gln Thr Gln Thr Ser Pro Glu His Leu Ala  
 245 250 255

Pro Gln Gln Asp Gln Val Pro Thr Gln Ala Gln Ser Gln Glu Gln Thr  
 260 265 270

Ser Glu Lys Thr Gln Asp Gln Pro Gln Thr Trp Pro Gln Gly Ser Val  
 275 280 285

Pro Pro Pro Glu Gln Ala Ser Gly Pro Ala Cys Ala Thr Glu Pro Gln  
 290 295 300

Leu Ser Ser His Ala Ala Glu Ala Gly Ser Asp Pro Asp Lys Ala Leu  
 305 310 315 320

Pro Glu Pro Val Ser Ala Gln Ser Ser Glu Asp Arg Ser Arg Glu Ala  
 325 330 335

Ser Ala Gly Gly Leu Asp Leu Gly Glu Cys Glu Lys Arg Ala Gly Glu  
 340 345 350

Met Leu Gly Met Trp Gly Ala Gly Ser Ser Leu Lys Val Thr Ile Leu

355		360		365
Gln Ser Ser Asn Ser Arg Ala Phe Asn Thr Thr Pro Leu Thr Ser Gly				
370		375		380
Pro Arg Pro Gly Asp Ser Thr Ser Ala Thr Pro Ala Ile Ala Ser Thr				
385		390		395 400
Pro Ser Lys Gln Ser Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser				
	405		410	415
Ser Ser Ser Gln Gln Glu Phe Gln Asp His Met Ser Glu Ala Gln His				
	420		425	430
Gln Gln Arg Leu Gly Glu Ile Gln His Ser Ser Gln Thr Cys Leu Leu				
	435		440	445
Ser Leu Leu Pro Met Pro Arg Asp Ile Leu Glu Lys Glu Ala Glu Asp				
	450		455	460
Pro Pro Pro Lys Arg Trp Cys Asn Thr Cys Gln Val Tyr Tyr Val Gly				
465		470		475 480
Asp Leu Ile Gln His Arg Arg Thr Gln Glu His Lys Val Ala Lys Gln				
	485		490	495
Ser Leu Arg Pro Phe Cys Thr Ile Cys Asn Arg Tyr Phe Lys Thr Pro				
	500		505	510
Arg Lys Phe Val Glu His Val Lys Ser Gln Gly His Lys Asp Lys Ala				
	515		520	525
Gln Glu Leu Lys Thr Leu Glu Lys Glu Thr Gly Ser Pro Asp Glu Asp				
	530		535	540
His Phe Ile Thr Val Asp Ala Val Gly Cys Phe Glu Ser Gly Gln Glu				
545		550		555 560
Glu Asp Glu Asp Asp Asp Glu Glu Glu Glu Glu Glu Gly Glu Ile Glu				
	565		570	575
Ala Glu Glu Glu Phe Cys Lys Gln Val Lys Pro Arg Glu Thr Ser Ser				
	580		585	590

Glu Gln Gly Lys Gly Ser Glu Thr Tyr Asn Pro Asn Thr Ala Tyr Gly  
 595 600 605

Glu Asp Phe Leu Val Pro Val Met Gly Tyr Val Cys Gln Ile Cys His  
 610 615 620

Lys Phe Tyr Asp Ser Asn Ser Glu Leu Arg Leu Ser His Cys Lys Ser  
 625 630 635 640

Leu Ala His Phe Glu Asn Leu Gln Lys Tyr Lys Ala Lys Asn Pro Ser  
 645 650 655

Pro Pro Pro Thr Arg Pro Val Ser Arg Lys Cys Ala Ile Asn Ala Arg  
 660 665 670

Asn Ala Leu Thr Ala Leu Phe Thr Ser Ser His Gln Pro Ser Pro Gln  
 675 680 685

Asp Thr Val Lys Met Pro Ser Lys Val Lys Pro Gly Ser Pro Gly Leu  
 690 695 700

Pro Pro Pro Leu Arg Arg Ser Thr Arg Leu Lys Thr  
 705 710 715

<210> 28  
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 <212> PRT  
 <213> Mouse

<400> 28

Met Phe Asn Pro Gln Leu Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln  
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Gln Gln Leu Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln  
 20 25 30

Gln Gln Ile Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln  
 35 40 45

Ala Ser Leu Ser Ile Pro Val Ser Arg Gly Leu Pro Gln Gln Ser Ser  
 50 55 60

Pro Gln Gln Leu Leu Ser Leu Gln Gly Leu His Ser Thr Ser Leu Leu  
65 70 75 80

Asn Gly Pro Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly  
85 90 95

Leu Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Gly Ala Ser Leu  
100 105 110

Thr Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Ala Phe Asn Val Thr  
115 120 125

Ala Pro Ser Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Met Val Thr  
130 135 140

Pro Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu  
145 150 155 160

Gly Pro Pro Pro Val Gly Val Pro Ile Asn Pro Ser Gln Leu Asn His  
165 170 175

Ser Gly Arg Asn Thr Gln Lys Gln Ala Arg Thr Pro Ser Ser Thr Thr  
180 185 190

Pro Asn Arg Lys Thr Val Pro Leu Glu Asp Arg Glu Asp Pro Thr Glu  
195 200 205

Gly Ser Glu Glu Ala Thr Glu Leu Gln Met Asp Thr Cys Glu Asp Gln  
210 215 220

Asp Ser Leu Val Gly Pro Asp Ser Met Leu Ser Glu Pro Gln Val Pro  
225 230 235 240

Glu Pro Glu Pro Phe Glu Thr Leu Glu Pro Pro Ala Lys Arg Cys Arg  
245 250 255

Ser Ser Glu Glu Ser Thr Glu Lys Gly Pro Thr Gly Gln Pro Gln Ala  
260 265 270

Arg Val Gln Pro Gln Thr Gln Met Thr Ala Pro Lys Gln Thr Gln Thr  
275 280 285

Pro Asp Arg Leu Pro Glu Pro Pro Glu Val Gln Met Leu Pro Arg Ile

290		295		300
Gln Pro Gln Ala Leu Gln Ile Gln Thr Gln Pro Lys Leu Leu Arg Gln				
305		310		315 320
Ala Gln Thr Gln Thr Ser Pro Glu His Leu Ala Pro Gln Gln Asp Gln				
		325		330 335
Val Pro Thr Gln Ala Gln Ser Gln Glu Gln Thr Ser Glu Lys Thr Gln				
		340		345 350
Asp Gln Pro Gln Thr Trp Pro Gln Gly Ser Val Pro Pro Pro Glu Gln				
		355		360 365
Ala Ser Gly Pro Ala Cys Ala Thr Glu Pro Gln Leu Ser Ser His Ala				
		370		375 380
Ala Glu Ala Gly Ser Asp Pro Asp Lys Ala Leu Pro Glu Pro Val Ser				
		385		390 395 400
Ala Gln Ser Ser Glu Asp Arg Ser Arg Glu Ala Ser Ala Gly Gly Leu				
		405		410 415
Asp Leu Gly Glu Cys Glu Lys Arg Ala Gly Glu Met Leu Gly Met Trp				
		420		425 430
Gly Ala Gly Ser Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asn Ser				
		435		440 445
Arg Ala Phe Asn Thr Thr Pro Leu Thr Ser Gly Pro Ser Pro Gly Asp				
		450		455 460
Ser Thr Ser Ala Thr Pro Ala Ile Ala Ser Thr Pro Ser Lys Gln Ser				
		465		470 475 480
Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser Ser Ser Ser Gln Gln				
		485		490 495
Glu Phe Gln Asp His Met Ser Glu Ala Gln His Gln Gln Arg Leu Gly				
		500		505 510
Glu Ile Gln His Ser Ser Gln Thr Cys Leu Leu Ser Leu Leu Pro Met				
		515		520 525

Pro Arg Asp Ile Leu Glu Lys Glu Ala Glu Asp Pro Pro Pro Lys Arg  
 530 535 540

Trp Cys Asn Thr Cys Gln Val Tyr Tyr Val Gly Asp Leu Ile Gln His  
 545 550 555 560

Arg Arg Thr Gln Glu His Lys Val Ala Lys Gln Ser Leu Arg Pro Phe  
 565 570 575

Cys Thr Ile Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu  
 580 585 590

His Val Lys Ser Gln Gly His Lys Asp Lys Ala Gln Glu Leu Lys Thr  
 595 600 605

Leu Glu Lys Glu Thr Gly Ser Pro Asp Glu Asp His Phe Ile Thr Val  
 610 615 620

Glu Ala Val Gly Cys Phe Glu Ser Gly Gln Glu Glu Asp Glu Asp Asp  
 625 630 635 640

Asp Glu Glu Glu Glu Glu Glu Gly Glu Ile Glu Ala Glu Glu Glu Phe  
 645 650 655

Cys Lys Gln Val Lys Pro Arg Glu Thr Ser Ser Glu Gln Gly Lys Gly  
 660 665 670

Ser Glu Thr Tyr Asn Pro Asn Thr Ala Tyr Gly Glu Asp Phe Leu Val  
 675 680 685

Pro Val Met Gly Tyr Val Cys Gln Ile Cys His Lys Phe Tyr Asp Ser  
 690 695 700

Asn Ser Glu Leu Arg Leu Ser His Cys Lys  
 705 710

<210> 29  
 <211> 898  
 <212> PRT  
 <213> Homo sapiens

<400> 29



Met Phe Ser Gln Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln  
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 Leu Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln  
 20 25 30  
 Gln Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln  
 35 40 45  
 Ala Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro  
 50 55 60  
 Gln Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu  
 65 70 75 80  
 Asn Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly  
 85 90 95  
 Leu Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu  
 100 105 110  
 Thr Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala  
 115 120 125  
 Ser Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr  
 130 135 140  
 Pro Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu  
 145 150 155 160  
 Gly Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu  
 165 170 175  
 Ser Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr  
 180 185 190  
 Pro Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys  
 195 200 205  
 Ser Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp  
 210 215 220  
 Thr Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys

225		230		235		240
Glu Lys Arg Thr	Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu					
	245			250		255
Leu Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu						
	260			265		270
Pro Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr						
	275			280		285
Val Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu						
	290			295		300
Ala Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala						
305		310		315		320
Gln Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln						
	325			330		335
Val Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu						
	340			345		350
His Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu						
	355			360		365
Ala Glu Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His						
370		375		380		
Ser Gln Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu						
385		390		395		400
Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln Pro Pro						
	405			410		415
Arg Gln Val Gln Leu Gln Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr						
	420			425		430
Pro Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln Glu His						
	435			440		445
Pro Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His Glu Gln						
450		455		460		

Pro His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln Thr Pro  
 465 470 475 480

Val Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala Val Glu  
 485 490 495

Ala Gly Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly Thr Gln  
 500 505 510

Val Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu Asp Val  
 515 520 525

Gly Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp Gly Ala  
 530 535 540

Gly Gly Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser Arg Ala  
 545 550 555 560

Phe Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp Ser Val  
 565 570 575

Ser Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln  
 580 585 590

Phe Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe  
 595 600 605

Gln Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly Glu Ile  
 610 615 620

Gln His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro Val Pro Arg  
 625 630 635 640

Asp Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg Trp Cys  
 645 650 655

Asn Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His Arg Arg  
 660 665 670

Thr Gln Asp His Lys Ile Ala Lys Gln Ser Leu Arg Pro Phe Cys Thr  
 675 680 685

Val Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu His Val  
690 695 700

Lys Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser Leu Glu  
705 710 715 720

Lys Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val Asp Ala  
725 730 735

Val Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp Asp Glu Asp  
740 745 750

Glu Glu Glu Ile Glu Val Glu Glu Glu Leu Cys Lys Gln Val Arg Ser  
755 760 765

Arg Asp Ile Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr Tyr Ser Pro  
770 775 780

Asn Thr Ala Tyr Gly Val Asp Phe Leu Val Pro Val Met Gly Tyr Ile  
785 790 795 800

Cys Arg Ile Cys His Lys Phe Tyr His Ser Asn Ser Gly Ala Gln Leu  
805 810 815

Ser His Cys Lys Ser Leu Gly His Phe Glu Asn Leu Gln Lys Tyr Lys  
820 825 830

Ala Ala Lys Asn Pro Ser Pro Thr Thr Arg Pro Val Ser Arg Arg Cys  
835 840 845

Ala Ile Asn Ala Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser Ser Gly  
850 855 860

Arg Pro Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro Ser Lys Val  
865 870 875 880

Thr Ala Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser Thr Arg Leu  
885 890 895

Lys Thr

<210> 30  
 <211> 898  
 <212> PRT  
 <213> Homo sapiens

<400> 30

Met Phe Ser Gln Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln  
 1 5 10 15

Leu Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln  
 20 25 30

Gln Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln  
 35 40 45

Ala Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro  
 50 55 60

Gln Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu  
 65 70 75 80

Asn Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly  
 85 90 95

Leu Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu  
 100 105 110

Thr Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala  
 115 120 125

Ser Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr  
 130 135 140

Pro Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu  
 145 150 155 160

Gly Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu  
 165 170 175

Ser Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr  
 180 185 190

Pro Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys  
 195 200 205

Ser Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp  
 210 215 220

Thr Pro Glu Asp Gln Asp Leu Leu Pro Cys Pro Glu Asp Ile Ala Lys  
 225 230 235 240

Glu Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu  
 245 250 255

Leu Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu  
 260 265 270

Pro Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr  
 275 280 285

Val Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu  
 290 295 300

Ala Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala  
 305 310 315 320

Gln Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln  
 325 330 335

Val Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu  
 340 345 350

His Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu  
 355 360 365

Ala Glu Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His  
 370 375 380

Ser Gln Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu  
 385 390 395 400

Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln Pro Pro  
 405 410 415

Arg Gln Val Gln Leu Gln Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr  
 420 425 430

Pro Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln Glu His  
435 440 445

Pro Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His Glu Gln  
450 455 460

Pro His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln Thr Pro  
465 470 475 480

Val Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala Val Glu  
485 490 495

Ala Gly Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly Thr Gln  
500 505 510

Val Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu Asp Val  
515 520 525

Gly Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp Gly Ala  
530 535 540

Gly Gly Ser Leu Lys Val Thr Ile Leu Gln Gly Ser Asp Ser Arg Ala  
545 550 555 560

Phe Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp Ser Val  
565 570 575

Ser Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln  
580 585 590

Phe Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe  
595 600 605

Gln Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly Glu Ile  
610 615 620

Gln His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro Val Pro Arg  
625 630 635 640

Asp Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg Trp Cys  
645 650 655

Asn Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His Arg Arg  
660 665 670

Thr Gln Asp His Lys Ile Ala Lys Gln Ser Leu Arg Pro Phe Cys Thr  
675 680 685

Val Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu His Val  
690 695 700

Lys Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser Leu Glu  
705 710 715 720

Lys Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val Asp Ala  
725 730 735

Val Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp Asp Glu Asp  
740 745 750

Glu Glu Glu Ile Glu Val Glu Glu Glu Leu Cys Lys Gln Val Arg Ser  
755 760 765

Arg Asp Ile Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr Tyr Ser Pro  
770 775 780

Asn Thr Ala Tyr Gly Val Asp Phe Leu Val Pro Val Met Gly Tyr Ile  
785 790 795 800

Cys Arg Ile Cys His Lys Phe Tyr His Ser Asn Ser Gly Ala Gln Leu  
805 810 815

Ser His Cys Lys Ser Leu Gly His Phe Glu Asn Leu Gln Lys Tyr Lys  
820 825 830

Ala Ala Lys Asn Pro Ser Pro Thr Thr Arg Pro Val Ser Arg Arg Cys  
835 840 845

Ala Ile Asn Ala Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser Ser Gly  
850 855 860

Arg Pro Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro Ser Lys Val  
865 870 875 880

Thr Ala Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser Thr Arg Leu



885

890

895

Lys Thr

<210> 31  
 <211> 896  
 <212> PRT  
 <213> Homo sapiens

&lt;400&gt; 31

Phe Ser Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln Leu Gln  
 1 5 10 15

Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln  
 20 25 30

Ser Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln Ala Pro  
 35 40 45

Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln Gln  
 50 55 60

Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn Gly  
 65 70 75 80

Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Leu Asp  
 85 90 95

Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu Thr Met  
 100 105 110

Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala Ser Pro  
 115 120 125

Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr Pro Asn  
 130 135 140

Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu Gly Pro  
 145 150 155 160

Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser Gly  
 165 170 175

Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro Asn  
180 185 190

Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys Ser Asp  
195 200 205

Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr Pro  
210 215 220

Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu Lys  
225 230 235 240

Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu Pro  
245 250 255

Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro Pro ...  
260 265 270

Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val Pro  
275 280 285

Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala Gln  
290 295 300

Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala Gln Val  
305 310 315 320

Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln Val Gln  
325 330 335

Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu His Leu  
340 345 350

Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu Ala Glu  
355 360 365

Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln  
370 375 380

Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu Lys Gln  
385 390 395 400

Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln Pro Pro Arg Gln  
 405 410 415

Val Gln Leu Gln Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr Pro Gln  
 420 425 430

Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln Glu His Pro Pro  
 435 440 445

Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His Glu Gln Pro His  
 450 455 460

Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln Thr Pro Val Val  
 465 470 475 480

Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala Val Glu Ala Gly  
 485 490 495

Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly Thr Gln Val Ser  
 500 505 510

Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu Asp Val Gly Glu  
 515 520 525

Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp Gly Ala Gly Gly  
 530 535 540

Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser Arg Ala Phe Ser  
 545 550 555 560

Thr Val Pro Leu Thr Leu Val Pro Arg Pro Ser Asp Ser Val Ser Ser  
 565 570 575

Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln Phe Phe  
 580 585 590

Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe Gln Asp  
 595 600 605

His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly Glu Ile Gln His  
 610 615 620

Met Ser Gln Ala Cys Leu Leu Pro Leu Leu Pro Val Pro Arg Asp Val

625						630						635				640
Leu	Glu	Thr	Glu	Asp	Glu	Glu	Pro	Pro	Pro	Arg	Arg	Trp	Cys	Asn	Thr	
				645					650					655		
Cys	Gln	Leu	Tyr	Tyr	Met	Gly	Asp	Leu	Ile	Gln	His	Arg	Arg	Thr	Gln	
			660					665					670			
Asp	His	Lys	Ile	Ala	Lys	Gln	Ser	Leu	Arg	Pro	Phe	Cys	Thr	Val	Cys	
		675					680					685				
Asn	Arg	Tyr	Phe	Lys	Thr	Pro	Arg	Lys	Phe	Val	Glu	His	Val	Lys	Ser	
	690					695					700					
Gln	Gly	His	Lys	Asp	Lys	Ala	Lys	Glu	Leu	Lys	Ser	Leu	Glu	Lys	Glu	
705					710					715					720	
Ile	Ala	Gly	Gln	Asp	Glu	Asp	His	Phe	Ile	Thr	Val	Gly	Ala	Val	Gly	
				725					730					735		
Cys	Phe	Glu	Gly	Asp	Glu	Glu	Glu	Glu	Glu	Asp	Asp	Glu	Asp	Glu	Glu	
			740					745					750			
Glu	Ile	Glu	Val	Glu	Glu	Glu	Leu	Cys	Lys	Gln	Val	Arg	Ser	Arg	Asp	
	755						760					765				
Ile	Ser	Arg	Glu	Glu	Trp	Lys	Gly	Ser	Glu	Thr	Tyr	Ser	Pro	Asn	Thr	
	770					775					780					
Ala	Tyr	Gly	Val	Asp	Phe	Leu	Val	Pro	Val	Met	Gly	Tyr	Ile	Cys	Arg	
785					790					795					800	
Ile	Cys	His	Lys	Phe	Tyr	His	Ser	Asn	Ser	Gly	Ala	Gln	Leu	Ser	His	
				805					810					815		
Cys	Lys	Ser	Leu	Gly	His	Phe	Glu	Asn	Leu	Gln	Lys	Tyr	Lys	Ala	Ala	
			820					825					830			
Lys	Asn	Pro	Ser	Pro	Thr	Thr	Arg	Pro	Val	Ser	Arg	Arg	Cys	Ala	Ile	
		835					840					845				
Asn	Ala	Arg	Asn	Ala	Leu	Thr	Ala	Leu	Phe	Thr	Ser	Ser	Gly	Arg	Pro	
	850					855					860					

Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro Ser Lys Val Thr Ala  
 865 870 875 880

Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser Thr Arg Leu Lys Thr  
 885 890 895

<210> 32  
 <211> 842  
 <212> PRT  
 <213> Homo sapiens

<400> 32

Met Phe Ser Gln Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln  
 1 5 10 15

Leu Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln  
 20 25 30

Gln Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln  
 35 40 45

Ala Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro  
 50 55 60

Gln Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu  
 65 70 75 80

Asn Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly  
 85 90 95

Leu Asp Gln Phe Val Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu  
 100 105 110

Thr Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala  
 115 120 125

Ser Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr  
 130 135 140

Pro Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu  
 145 150 155 160

Gly Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu  
 165 170 175

Ser Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr  
 180 185 190

Pro Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys  
 195 200 205

Ser Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp  
 210 215 220

Thr Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys  
 225 230 235 240

Glu Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu  
 245 250 255

Leu Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu  
 260 265 270

Pro Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr  
 275 280 285

Val Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu  
 290 295 300

Ala Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala  
 305 310 315 320

Gln Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln  
 325 330 335

Val Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu  
 340 345 350

His Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu  
 355 360 365

Ala Glu Pro Gln Lys Gln Val Gln Pro Gln Val His Thr Gln Ala Gln  
 370 375 380

Pro Ser Val Gln Pro Gln Glu His Pro Pro Ala Gln Val Ser Val Gln

385		390		395		400									
Pro	Pro	Glu	Gln	Thr	His	Glu	Gln	Pro	His	Thr	Gln	Pro	Gln	Val	Ser
				405					410					415	
Leu	Leu	Ala	Pro	Glu	Gln	Thr	Pro	Val	Val	Val	His	Val	Cys	Gly	Leu
			420					425					430		
Glu	Met	Pro	Pro	Asp	Ala	Val	Glu	Ala	Gly	Gly	Gly	Met	Glu	Lys	Thr
		435					440					445			
Leu	Pro	Glu	Pro	Val	Gly	Thr	Gln	Val	Ser	Met	Glu	Glu	Ile	Gln	Asn
	450					455					460				
Glu	Ser	Ala	Cys	Gly	Leu	Asp	Val	Gly	Glu	Cys	Glu	Asn	Arg	Ala	Arg
465					470					475					480
Glu	Met	Pro	Gly	Val	Trp	Gly	Ala	Gly	Gly	Ser	Leu	Lys	Val	Thr	Ile
			485					490						495	
Leu	Gln	Ser	Ser	Asp	Ser	Arg	Ala	Phe	Ser	Thr	Val	Pro	Leu	Thr	Pro
			500					505					510		
Val	Pro	Arg	Pro	Ser	Asp	Ser	Val	Ser	Ser	Thr	Pro	Ala	Ala	Thr	Ser
		515					520					525			
Thr	Pro	Ser	Lys	Gln	Ala	Leu	Gln	Phe	Phe	Cys	Tyr	Ile	Cys	Lys	Ala
	530					535					540				
Ser	Cys	Ser	Ser	Gln	Gln	Glu	Phe	Gln	Asp	His	Met	Ser	Glu	Pro	Gln
545				550						555					560
His	Gln	Gln	Arg	Leu	Gly	Glu	Ile	Gln	His	Met	Ser	Gln	Ala	Cys	Leu
			565					570						575	
Leu	Ser	Leu	Leu	Pro	Met	Pro	Arg	Asp	Val	Leu	Glu	Thr	Glu	Asp	Glu
			580					585					590		
Glu	Pro	Pro	Pro	Arg	Arg	Trp	Cys	Asn	Thr	Cys	Gln	Leu	Tyr	Tyr	Met
		595					600					605			
Gly	Asp	Leu	Ile	Gln	His	Arg	Arg	Thr	Gln	Asp	His	Lys	Val	Ala	Lys
	610					615					620				

Gln Pro Leu Arg Pro Phe Cys Thr Val Cys Asn Arg Tyr Phe Lys Thr  
 625 630 635 640

Pro Arg Lys Phe Val Glu His Val Lys Ser Gln Gly His Lys Asp Lys  
 645 650 655

Ala Lys Glu Leu Lys Ser Leu Glu Lys Glu Ile Ala Gly Gln Asp Glu  
 660 665 670

Asp His Phe Ile Thr Val Asp Ala Val Gly Cys Phe Glu Gly Asp Glu  
 675 680 685

Glu Glu Glu Glu Asp Asp Glu Asp Glu Glu Glu Ile Lys Val Glu Glu  
 690 695 700

Glu Leu Cys Lys Gln Val Arg Ser Arg Asp Ile Ser Arg Glu Glu Trp  
 705 710 715 720

Lys Gly Ser Glu Thr Tyr Ser Pro Asn Thr Ala Tyr Gly Val Asp Phe  
 725 730 735

Leu Val Pro Val Met Gly Tyr Ile Cys Arg Ile Cys His Lys Phe Tyr  
 740 745 750

His Ser Asn Ser Gly Ala Gln Leu Ser His Cys Lys Ser Leu Gly His  
 755 760 765

Phe Glu Asn Leu Gln Lys Tyr Lys Ala Ala Lys Asn Pro Ser Pro Thr  
 770 775 780

Thr Arg Pro Val Ser Arg Arg Cys Ala Ile Asn Ala Arg Asn Ala Leu  
 785 790 795 800

Thr Ala Leu Phe Thr Ser Ser Gly Arg Pro Pro Ser Gln Pro Asn Thr  
 805 810 815

Gln Asp Lys Thr Pro Ser Lys Val Thr Ala Arg Pro Ser Gln Pro Pro  
 820 825 830

Leu Pro Arg Arg Ser Thr Arg Leu Lys Thr  
 835 840



<210> 33  
 <211> 837  
 <212> PRT  
 <213> Homo sapiens

<400> 33

Met Phe Ser Gln Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Ala  
 1 5 10 15

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
 20 25 30

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
 35 40 45

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Leu  
 50 55 60

Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu Thr  
 65 70 75 80

Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala Ser  
 85 90 95

Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr Pro  
 100 105 110

Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu Gly  
 115 120 125

Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser  
 130 135 140

Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro  
 145 150 155 160

Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys Ser  
 165 170 175

Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr  
 180 185 190

Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu

195		200		205
Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu				
210		215		220
Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro				
225		230		235 240
Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val				
	245		250	255
Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala				
	260		265	270
Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala Gln				
	275		280	285
Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln Val				
	290		295	300
Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu His				
305		310		315 320
Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu Ala				
	325		330	335
Glu Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser				
	340		345	350
Gln Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu Lys				
	355		360	365
Gln Val Gln Pro Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro				
	370		375	380
Gln Glu His Pro Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr				
385		390		395 400
His Glu Gln Pro His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu				
	405		410	415
Gln Thr Pro Val Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp				
	420		425	430

Ala Val Glu Ala Gly Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val  
 435 440 445

Gly Thr Gln Val Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly  
 450 455 460

Leu Asp Val Gly Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val  
 465 470 475 480

Trp Gly Ala Gly Gly Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp  
 485 490 495

Ser Arg Ala Phe Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser  
 500 505 510

Asp Ser Val Ser Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln  
 515 520 525

Ala Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln  
 530 535 540

Gln Glu Phe Gln Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu  
 545 550 555 560

Gly Glu Ile Gln His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro  
 565 570 575

Val Pro Arg Asp Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg  
 580 585 590

Arg Trp Cys Asn Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln  
 595 600 605

His Arg Arg Thr Gln Asp His Lys Ile Ala Lys Gln Ser Leu Arg Pro  
 610 615 620

Phe Cys Thr Val Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val  
 625 630 635 640

Glu His Val Lys Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys  
 645 650 655

Ser Leu Glu Lys Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr  
660 665 670

Val Asp Ala Val Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp  
675 680 685

Asp Glu Asp Glu Glu Glu Ile Glu Val Glu Glu Glu Leu Cys Lys Gln  
690 695 700

Val Arg Ser Arg Asp Ile Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr  
705 710 715 720

Tyr Ser Pro Asn Thr Ala Tyr Gly Val Asp Phe Leu Val Pro Val Met  
725 730 735

Gly Tyr Ile Cys Arg Ile Cys His Lys Phe Tyr His Ser Asn Ser Gly  
740 745 750

Ala Gln Leu Ser His Cys Lys Ser Leu Gly His Phe Glu Asn Leu Gln  
755 760 765

Lys Tyr Lys Ala Ala Lys Asn Pro Ser Pro Thr Thr Arg Pro Val Ser  
770 775 780

Arg Arg Cys Ala Ile Asn Ala Arg Asn Ala Leu Thr Ala Leu Phe Thr  
785 790 795 800

Ser Ser Gly Arg Pro Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro  
805 810 815

Ser Lys Val Thr Ala Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser  
820 825 830

Thr Arg Leu Lys Thr  
835

<210> 34  
<211> 818  
<212> PRT  
<213> Homo sapiens

<400> 34

Met Phe Ser Gln Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln

1	5	10	15
Leu Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln	20	25	30
Gln Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln	35	40	45
Ala Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro	50	55	60
Gln Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu	65	70	80
Asn Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly	85	90	95
Asn Leu Arg Gly Tyr Gly Met Ala Ser Pro Gly Leu Ala Ala Pro Ser	100	105	110
Leu Thr Pro Pro Gln Leu Ala Thr Pro Asn Leu Gln Gln Phe Phe Pro	115	120	125
Gln Ala Thr Arg Gln Ser Leu Leu Gly Pro Pro Pro Val Gly Val Pro	130	135	140
Met Asn Pro Ser Gln Phe Asn Leu Ser Gly Arg Asn Pro Gln Lys Gln	145	150	160
Ala Arg Thr Ser Ser Ser Thr Thr Pro Asn Arg Lys Asp Ser Ser Ser	165	170	175
Gln Thr Met Pro Val Glu Asp Lys Ser Asp Pro Pro Glu Gly Ser Glu	180	185	190
Glu Ala Ala Glu Pro Arg Met Asp Thr Pro Glu Asp Gln Asp Leu Pro	195	200	205
Pro Cys Pro Glu Asp Ile Ala Lys Glu Lys Arg Thr Pro Ala Pro Glu	210	215	220
Pro Glu Pro Cys Glu Ala Ser Glu Leu Pro Ala Lys Arg Leu Arg Ser	225	230	240

Ser Glu Glu Pro Thr Glu Lys Glu Pro Pro Gly Gln Leu Gln Val Lys  
245 250 255

Ala Gln Pro Gln Ala Arg Met Thr Val Pro Lys Gln Thr Gln Thr Pro  
260 265 270

Asp Leu Leu Pro Glu Ala Leu Glu Ala Gln Val Leu Pro Arg Phe Gln  
275 280 285

Pro Arg Val Leu Gln Val Gln Ala Gln Val Gln Ser Gln Thr Gln Pro  
290 295 300

Arg Ile Pro Ser Thr Asp Thr Gln Val Gln Pro Lys Leu Gln Lys Gln  
305 310 315 320

Ala Gln Thr Gln Thr Ser Pro Glu His Leu Val Leu Gln Gln Lys Gln  
325 330 335

Val Gln Pro Gln Leu Gln Gln Glu Ala Glu Pro Gln Lys Gln Val Gln  
340 345 350

Pro Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln Glu His  
355 360 365

Pro Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His Glu Gln  
370 375 380

Pro His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln Thr Pro  
385 390 395 400

Val Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala Val Glu  
405 410 415

Ala Gly Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly Thr Gln  
420 425 430

Val Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu Asp Val  
435 440 445

Gly Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp Gly Ala  
450 455 460

Gly Gly Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser Arg Ala  
 465 470 475 480

Phe Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp Ser Val  
 485 490 495

Ser Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln  
 500 505 510

Phe Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe  
 515 520 525

Gln Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly Glu Ile  
 530 535 540

Gln His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro Val Pro Arg  
 545 550 555 560

Asp Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg Trp Cys  
 565 570 575

Asn Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His Arg Arg  
 580 585 590

Thr Gln Asp His Lys Ile Ala Lys Gln Ser Leu Arg Pro Phe Cys Thr  
 595 600 605

Val Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu His Val  
 610 615 620

Lys Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser Leu Glu  
 625 630 635 640

Lys Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val Asp Ala  
 645 650 655

Val Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp Asp Glu Asp  
 660 665 670

Glu Glu Glu Ile Glu Val Glu Glu Glu Leu Cys Lys Gln Val Arg Ser  
 675 680 685

Arg Asp Ile Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr Tyr Ser Pro  
690 695 700

Asn Thr Ala Tyr Gly Val Asp Phe Leu Val Pro Val Met Gly Tyr Ile  
705 710 715 720

Cys Arg Ile Cys His Lys Phe Tyr His Ser Asn Ser Gly Ala Gln Leu  
725 730 735

Ser His Cys Lys Ser Leu Gly His Phe Glu Asn Leu Gln Lys Tyr Lys  
740 745 750

Ala Ala Lys Asn Pro Ser Pro Thr Thr Arg Pro Val Ser Arg Arg Cys  
755 760 765

Ala Ile Asn Ala Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser Ser Gly  
770 775 780

Arg Pro Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro Ser Lys Val  
785 790 795 800

Thr Ala Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser Thr Arg Leu  
805 810 815

Lys Thr

<210> 35  
<211> 820  
<212> PRT  
<213> Homo sapiens

<400> 35

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
1 5 10 15

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
20 25 30

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Asn  
35 40 45

Leu Arg Gly Tyr Gly Met Ala Ser Pro Gly Leu Ala Ala Pro Ser Leu  
50 55 60



Thr Pro Pro Gln Leu Ala Thr Pro Asn Leu Gln Gln Phe Phe Pro Gln  
65 70 75 80

Ala Thr Arg Gln Ser Leu Leu Gly Pro Pro Pro Val Gly Val Pro Met  
85 90 95

Asn Pro Ser Gln Phe Asn Leu Ser Gly Arg Asn Pro Gln Lys Gln Ala  
100 105 110

Arg Thr Ser Ser Ser Thr Thr Pro Asn Arg Lys Thr Met Pro Val Glu  
115 120 125

Asp Lys Ser Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg  
130 135 140

Met Asp Thr Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile  
145 150 155 160

Ala Lys Glu Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala  
165 170 175

Ser Glu Leu Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu  
180 185 190

Lys Glu Pro Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg  
195 200 205

Met Thr Val Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala  
210 215 220

Leu Glu Ala Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val  
225 230 235 240

Gln Ala Gln Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp  
245 250 255

Thr Gln Val Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser  
260 265 270

Pro Glu His Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln  
275 280 285

Gln Glu Ala Glu Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln  
 290 295 300

Ala His Ser Gln Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu  
 305 310 315 320

Pro Leu Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln  
 325 330 335

Pro Pro Arg Gln Val Gln Leu Gln Leu Gln Lys Gln Val Gln Thr Gln  
 340 345 350

Thr Tyr Pro Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln  
 355 360 365

Glu His Pro Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His  
 370 375 380

Glu Gln Pro His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln  
 385 390 395 400

Thr Pro Val Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala  
 405 410 415

Val Glu Ala Gly Gly Ser Met Glu Lys Thr Leu Pro Glu Pro Val Gly  
 420 425 430

Thr Gln Val Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu  
 435 440 445

Asp Val Gly Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp  
 450 455 460

Gly Ala Gly Gly Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser  
 465 470 475 480

Arg Ala Phe Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp  
 485 490 495

Ser Val Ser Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala  
 500 505 510

Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln  
 515 520 525

Glu Phe Gln Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly  
 530 535 540

Glu Ile Gln His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro Val  
 545 550 555 560

Pro Arg Asp Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg  
 565 570 575

Trp Cys Asn Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His  
 580 585 590

Arg Arg Thr Gln Asp His Arg Ile Ala Lys Gln Ser Leu Arg Pro Phe  
 595 600 605

Cys Thr Val Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu  
 610 615 620

His Val Lys Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser  
 625 630 635 640

Leu Glu Lys Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val  
 645 650 655

Asp Ala Val Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp Asp  
 660 665 670

Glu Asp Glu Glu Glu Ile Glu Val Glu Glu Glu Leu Cys Lys Gln Val  
 675 680 685

Arg Ser Arg Asp Ile Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr Tyr  
 690 695 700

Ser Pro Asn Thr Ala Tyr Gly Val Asp Phe Leu Val Pro Val Met Gly  
 705 710 715 720

Tyr Ile Cys Arg Ile Cys His Lys Phe Tyr His Asn Asn Ser Gly Ala  
 725 730 735

Gln Leu Ser His Cys Lys Ser Leu Gly His Phe Glu Asn Leu Gln Lys

740	745	750
Tyr Lys Ala Ala Lys Asn Pro Ser Pro Thr Thr Arg Pro Val Ser Arg		
755	760	765
Arg Cys Ala Ile Asn Ala Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser		
770	775	780
Ser Gly Arg Pro Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro Ser		
785	790	795
Lys Val Thr Ala Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser Thr		
805	810	815
Arg Leu Lys Thr		
820		
<210> 36		
<211> 414		
<212> PRT		
<213> Homo sapiens		
<400> 36		
Met Phe Ser Gln Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln		
1	5	10
Leu Gln Gln Leu Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln		
20	25	30
Gln Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln		
35	40	45
Ala Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro		
50	55	60
Gln Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu		
65	70	75
Asn Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly		
85	90	95
Asn Leu Arg Gly Tyr Gly Met Ala Ser Pro Gly Leu Ala Ala Pro Ser		
100	105	110

Leu Thr Pro Pro Gln Leu Ala Thr Pro Asn Leu Gln Gln Phe Phe Pro  
115 120 125

Gln Ala Thr Arg Gln Ser Leu Leu Gly Pro Pro Pro Val Gly Val Pro  
130 135 140

Met Asn Pro Ser Gln Phe Asn Leu Ser Gly Arg Asn Pro Gln Lys Gln  
145 150 155 160

Ala Arg Thr Ser Ser Ser Thr Thr Pro Asn Arg Lys Asp Ser Ser Ser  
165 170 175

Gln Thr Met Pro Val Glu Asp Lys Ser Asp Pro Pro Glu Gly Ser Glu  
180 185 190

Glu Ala Ala Glu Pro Arg Met Asp Thr Pro Glu Asp Gln Asp Leu Pro  
195 200 205

Pro Cys Pro Glu Asp Ile Ala Lys Glu Lys Arg Thr Pro Ala Pro Glu  
210 215 220

Pro Glu Pro Cys Glu Ala Ser Glu Leu Pro Ala Lys Arg Leu Arg Ser  
225 230 235 240

Ser Glu Glu Pro Thr Glu Lys Glu Pro Pro Gly Gln Leu Gln Val Lys  
245 250 255

Ala Gln Pro Gln Ala Arg Met Thr Val Pro Lys Gln Thr Gln Thr Pro  
260 265 270

Asp Leu Leu Pro Glu Ala Leu Glu Ala Gln Val Leu Pro Arg Phe Gln  
275 280 285

Pro Arg Val Leu Gln Val Gln Ala Gln Val Gln Ser Gln Thr Gln Pro  
290 295 300

Arg Ile Pro Ser Thr Asp Thr Gln Val Gln Pro Lys Leu Gln Lys Gln  
305 310 315 320

Ala Gln Thr Gln Thr Ser Pro Glu His Leu Val Leu Gln Gln Lys Gln  
325 330 335

Val Gln Pro Gln Leu Gln Gln Glu Ala Glu Pro Gln Lys Gln Val Gln  
 340 345 350

Pro Gln Val Gln Pro Gln Ala His Ser Gln Gly Pro Arg Gln Val Gln  
 355 360 365

Leu Gln Gln Glu Ala Glu Pro Leu Lys Gln Val Gln Pro Gln Val Gln  
 370 375 380

Pro Gln Ala His Ser Gln Pro His Leu Pro Gln Val Leu Ser Gln Gln  
 385 390 395 400

Leu Arg Gly Thr Ala Leu Pro Leu Gln Val Pro Gly Pro Leu  
 405 410

<210> 37  
 <211> 75  
 <212> PRT  
 <213> Homo sapiens

<400> 37

Leu Gln Gln Gln Gln Gln Gln Leu Gln Gln Leu Gln Gln Gln Gln Leu  
 1 5 10 15

Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Leu Gln Leu Gln Gln Leu  
 20 25 30

Leu Gln Gln Ser Pro Pro Gln Ala Pro Leu Pro Met Ala Val Ser Arg  
 35 40 45

Gly Leu Pro Pro Gln Gln Pro Gln Gln Pro Leu Leu Asn Leu Gln Gly  
 50 55 60

Thr Asn Ser Ala Ser Leu Leu Asn Gly Ser Met  
 65 70 75

<210> 38  
 <211> 33  
 <212> PRT  
 <213> Homo sapiens

<400> 38

Gln Gln Leu Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu  
 1 5 10 15

Gln Gln Gln Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro  
 20 25 30

Pro

<210> 39  
 <211> 52  
 <212> PRT  
 <213> Homo sapiens

<400> 39

Met Phe Ser Gln Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln  
 1 5 10 15

Leu Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln  
 20 25 30

Gln Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln  
 35 40 45

Ala Pro Leu Pro  
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<210> 40  
 <211> 26  
 <212> PRT  
 <213> Homo sapiens

<400> 40

Pro Pro Thr Pro Arg Arg Asp Val Phe Ala His Val Pro Val Gln Gly  
 1 5 10 15

Trp Ser Thr Ala Arg Leu Val Thr Asp Met  
 20 25

<210> 41  
 <211> 24  
 <212> PRT  
 <213> Homo sapiens

<400> 41

Gly Leu Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly  
 1 5 10 15

Leu Thr Met Pro Thr Ala Thr Leu  
20

<210> 42  
<211> 56  
<212> PRT  
<213> Homo sapiens

<400> 42

Pro Gln Val Gln Pro Gln Ala His Ser Gln Gly Pro Arg Gln Val Gln  
1 5 10 15

Leu Gln Gln Glu Ala Glu Pro Leu Lys Gln Val Gln Pro Gln Val Gln  
20 25 30

Pro Gln Ala His Ser Gln Pro Pro Arg Gln Val Gln Leu Gln Leu Gln  
35 40 45

Lys Gln Val Gln Thr Gln Thr Tyr  
50 55

<210> 43  
<211> 28  
<212> PRT  
<213> Homo sapiens

<400> 43

Pro Gln Val Gln Pro Gln Ala His Ser Gln Pro Pro Arg Gln Val Gln  
1 5 10 15

Leu Gln Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr  
20 25

<210> 44  
<211> 112  
<212> PRT  
<213> Homo sapiens

<400> 44

Gln Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln  
1 5 10 15

Val Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu  
20 25 30



His Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu  
 35 40 45

Ala Glu Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His  
 50 55 60

Ser Gln Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu  
 65 70 75 80

Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln Pro Pro  
 85 90 95

Arg Gln Val Gln Leu Gln Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr  
 100 105 110

<210> 45  
 <211> 2687  
 <212> DNA  
 <213> Mouse

<400> 45  
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atccaataaa gagtcagtag tttggcaaaa aaaaaaaaaa aaaaaaa 2687

<210> 46  
 <211> 2922  
 <212> DNA  
 <213> Homo sapiens

<400> 46  
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ggtagtttgg ctgtgcaaaa aaaaaaaaaa aaaaaaaaaa aa 2922

<210> 47  
<211> 897  
<212> PRT  
<213> Homo sapiens

<400> 47

Met Phe Ser Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Leu  
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Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln  
20 25 30

Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln Ala  
35 40 45

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
50 55 60

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
65 70 75 80

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Leu  
85 90 95

Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu Thr  
100 105 110

Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala Ser  
115 120 125

Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr Pro  
130 135 140

Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu Gly  
145 150 155 160

Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser  
165 170 175

Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro  
180 185 190

Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys Ser  
195 200 205

Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr  
210 215 220

Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu  
 225 230 235 240

Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu  
 245 250 255

Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro  
 260 265 270

Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val  
 275 280 285

Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala  
 290 295 300

Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala Gln  
 305 310 315 320

Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln Val  
 325 330 335

Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu His  
 340 345 350

Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu Ala  
 355 360 365

Glu Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser  
 370 375 380

Gln Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu Lys  
 385 390 395 400

Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln Pro Pro Arg  
 405 410 415

Gln Val Gln Leu Gln Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr Pro  
 420 425 430

Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln Glu His Pro  
 435 440 445

Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His Glu Gln Pro  
 450 455 460

His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln Thr Pro Val  
 465 470 475 480

Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala Val Glu Ala  
 485 490 495

Gly Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly Thr Gln Val  
 500 505 510

Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu Asp Val Gly  
 515 520 525

Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp Gly Ala Gly  
 530 535 540

Gly Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser Arg Ala Phe  
 545 550 555 560

Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp Ser Val Ser  
 565 570 575

Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln Phe  
 580 585 590

Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe Gln  
 595 600 605

Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly Glu Ile Gln  
 610 615 620

His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro Val Pro Arg Asp  
 625 630 635 640

Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg Trp Cys Asn  
 645 650 655

Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His Arg Arg Thr  
 660 665 670

Gln Asp His Lys Ile Ala Lys Gln Ser Leu Arg Pro Phe Cys Thr Val

675		680		685
Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu His Val Lys				
690		695		700
Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser Leu Glu Lys				
705		710		715 720
Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val Asp Ala Val				
		725		730 735
Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp Asp Glu Asp Glu				
		740		745 750
Glu Glu Ile Glu Val Glu Glu Glu Leu Cys Lys Gln Val Arg Ser Arg				
		755		760 765
Asp Ile Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr Tyr Ser Pro Asn				
		770		775 780
Thr Ala Tyr Gly Val Asp Phe Leu Val Pro Val Met Gly Tyr Ile Cys				
		785		790 795 800
Arg Ile Cys His Lys Phe Tyr His Ser Asn Ser Gly Ala Gln Leu Ser				
		805		810 815
His Cys Lys Ser Leu Gly His Phe Glu Asn Leu Gln Lys Tyr Lys Ala				
		820		825 830
Ala Lys Asn Pro Ser Pro Thr Thr Arg Pro Val Ser Arg Arg Cys Ala				
		835		840 845
Ile Asn Ala Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser Ser Gly Arg				
		850		855 860
Pro Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro Ser Lys Val Thr				
		865		870 875 880
Ala Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser Thr Arg Leu Lys				
		885		890 895

Thr



<210> 48  
 <211> 49  
 <212> PRT  
 <213> Homo sapiens

<400> 48

Met Phe Ser Gln Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln  
 1 5 10 15

Leu Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln  
 20 25 30

Gln Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln  
 35 40 45

Ala

<210> 49  
 <211> 215  
 <212> DNA  
 <213> Homo sapiens

<400> 49

tgggggctgc gggggccggcc catccgtggg ggcgacttga gcgttgaggg cgcgcgggga 60  
 ggcgagccac catgttcagc cagcagcagc agcagctcca gcaacagcag cagcagctcc 120  
 agcagttaca gcagcagcag ctccagcagc agcaattgca gcagcagcag ttactgcagc 180  
 tccagcagct gctccagcag tccccaccac aggcc 215

<210> 50  
 <211> 101  
 <212> DNA  
 <213> Homo sapiens

<400> 50

cagcagctcc agcagttaca gcagcagcag ctccagcagc agcaattgca gcagcagcag 60  
 ttactgcagc tccagcagct gctccagcag tccccaccac a 101

<210> 51  
 <211> 72  
 <212> DNA  
 <213> Homo sapiens

<400> 51

ggactggacc agtttgcaat gccaccagcc acgtatgaca ctgccggtct caccatgccc 60  
 acagcaacac tg 72  
  
 <210> 52  
 <211> 15  
 <212> DNA  
 <213> Homo sapiens  
  
 <400> 52  
 aggattcttc ttctc 15  
  
 <210> 53  
 <211> 86  
 <212> DNA  
 <213> Homo sapiens  
  
 <400> 53  
 ccacaggtgc agccccaggc acattcacag cccccaaggc aggtgcagct gcagctgcag 60  
 aagcaggtcc agacacagac atatcc 86  
  
 <210> 54  
 <211> 168  
 <212> DNA  
 <213> Homo sapiens  
  
 <400> 54  
 ccacaggtac agccacaggc acattcacag ggcccaaggc aggtgcagct gcagcaggag 60  
 gcagagccgc tgaagcaggt gcagccacag gtgcagcccc aggcacattc acagccccca 120  
 aggcaggtgc agctgcagct gcagaagcag gtccagacac agacatat 168  
  
 <210> 55  
 <211> 336  
 <212> DNA  
 <213> Homo sapiens  
  
 <400> 55  
 caggtgcagt cacagactca gccgcggata ccatccacag acaccaggt gcagccaaag 60  
 ctgcagaagc aggcgcaaac acagacctct ccagagcact tagtgctgca acagaagcag 120  
 gtgcagccac agctgcagca ggaggcagag ccacagaagc aggtgcagcc acaggtacag 180  
 ccacaggcac attcacaggg cccaaggcag gtgcagctgc agcaggaggc agagccgctg 240  
 aagcaggtgc agccacaggt gcagccccag gcacattcac agcccccaag gcaggtgcag 300  
 ctgcagctgc agaagcaggt ccagacacag acatat 336

<210> 56  
 <211> 24  
 <212> DNA  
 <213> Homo sapiens

<400> 56  
 gttgaggagg aactctgcaa gcag 24

<210> 57  
 <211> 78  
 <212> DNA  
 <213> Homo sapiens

<400> 57  
 gccacccaca ccacgaagag atgtgtttgc ccacgttcca gtgcaggggt ggagcacagc 60  
 ccggcttggt acagatat 78

<210> 58  
 <211> 863  
 <212> PRT  
 <213> Homo sapiens

<400> 58

Met	Phe	Ser	Gln	Gln	Gln	Gln	Gln	Leu	Gln	Gln	Gln	Gln	Gln	Ala	Pro
1			5					10						15	
Leu	Pro	Met	Ala	Val	Ser	Arg	Gly	Leu	Pro	Pro	Gln	Gln	Pro	Gln	Gln
			20					25					30		
Pro	Leu	Leu	Asn	Leu	Gln	Gly	Thr	Asn	Ser	Ala	Ser	Leu	Leu	Asn	Gly
			35				40					45			
Ser	Met	Leu	Gln	Arg	Ala	Leu	Leu	Leu	Gln	Gln	Leu	Gln	Gly	Leu	Asp
	50					55					60				
Gln	Phe	Ala	Met	Pro	Pro	Ala	Thr	Tyr	Asp	Thr	Ala	Gly	Leu	Thr	Met
65				70						75				80	
Pro	Thr	Ala	Thr	Leu	Gly	Asn	Leu	Arg	Gly	Tyr	Gly	Met	Ala	Ser	Pro
				85					90					95	
Gly	Leu	Ala	Ala	Pro	Ser	Leu	Thr	Pro	Pro	Gln	Leu	Ala	Thr	Pro	Asn
			100					105					110		
Leu	Gln	Gln	Phe	Phe	Pro	Gln	Ala	Thr	Arg	Gln	Ser	Leu	Leu	Gly	Pro
			115				120					125			

Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser Gly  
 130 135 140

Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro Asn  
 145 150 155 160

Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys Ser Asp  
 165 170 175

Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr Pro  
 180 185 190

Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu Lys  
 195 200 205

Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu Pro  
 210 215 220

Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro Pro  
 225 230 235 240

Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val Pro  
 245 250 255

Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala Gln  
 260 265 270

Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala Gln Val  
 275 280 285

Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln Val Gln  
 290 295 300

Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu His Leu  
 305 310 315 320

Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu Ala Glu  
 325 330 335

Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln  
 340 345 350

Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu Lys Gln  
 355 360 365

Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln Pro Pro Arg Gln  
 370 375 380

Val Gln Leu Gln Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr Pro Gln  
 385 390 395 400

Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln Glu His Pro Pro  
 405 410 415

Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His Glu Gln Pro His  
 420 425 430

Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln Thr Pro Val Val  
 435 440 445

Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala Val Glu Ala Gly  
 450 455 460

Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly Thr Gln Val Ser  
 465 470 475 480

Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu Asp Val Gly Glu  
 485 490 495

Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp Gly Ala Gly Gly  
 500 505 510

Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser Arg Ala Phe Ser  
 515 520 525

Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp Ser Val Ser Ser  
 530 535 540

Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln Phe Phe  
 545 550 555 560

Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe Gln Asp  
 565 570 575

His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly Glu Ile Gln His  
 580 585 590

Met Ser Gln Ala Leu Leu Ser Leu Leu Pro Val Pro Arg Asp Val Leu  
 595 600 605

Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg Trp Cys Asn Thr Cys  
 610 615 620

Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His Arg Arg Thr Gln Asp  
 625 630 635 640

His Lys Ile Ala Lys Gln Ser Leu Arg Pro Phe Cys Thr Val Cys Asn  
 645 650 655

Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu His Val Lys Ser Gln  
 660 665 670

Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser Leu Glu Lys Glu Ile  
 675 680 685

Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val Asp Ala Val Gly Cys  
 690 695 700

Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp Asp Glu Asp Glu Glu Glu  
 705 710 715 720

Ile Glu Val Glu Glu Glu Leu Cys Lys Gln Val Arg Ser Arg Asp Ile  
 725 730 735

Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr Tyr Ser Pro Asn Thr Ala  
 740 745 750

Tyr Gly Val Asp Phe Leu Val Pro Val Met Gly Tyr Ile Cys Arg Ile  
 755 760 765

Cys His Lys Phe Tyr His Ser Asn Ser Gly Ala Gln Leu Ser His Cys  
 770 775 780

Lys Ser Leu Gly His Phe Glu Asn Leu Gln Lys Tyr Lys Ala Ala Lys  
 785 790 795 800

Asn Pro Ser Pro Thr Thr Arg Pro Val Ser Arg Arg Cys Ala Ile Asn



Asn	Pro	Ser	Gln	Phe	Asn	Leu	Ser	Gly	Arg	Asn	Pro	Gln	Lys	Gln	Ala	
145					150					155					160	
Arg	Thr	Ser	Ser	Ser	Thr	Thr	Pro	Asn	Arg	Lys	Asp	Ser	Ser	Ser	Gln	
				165					170						175	
Thr	Met	Pro	Val	Glu	Asp	Lys	Ser	Asp	Pro	Pro	Glu	Gly	Ser	Glu	Glu	
			180					185					190			
Ala	Ala	Glu	Pro	Arg	Met	Asp	Thr	Pro	Glu	Asp	Gln	Asp	Leu	Pro	Pro	
		195					200					205				
Cys	Pro	Glu	Asp	Ile	Ala	Lys	Glu	Lys	Arg	Thr	Pro	Ala	Pro	Glu	Pro	
	210					215					220					
Glu	Pro	Cys	Glu	Ala	Ser	Glu	Leu	Pro	Ala	Lys	Arg	Leu	Arg	Ser	Ser	
225					230					235					240	
Glu	Glu	Pro	Thr	Glu	Lys	Glu	Pro	Pro	Gly	Gln	Leu	Gln	Val	Lys	Ala	
				245					250					255		
Gln	Pro	Gln	Ala	Arg	Met	Thr	Val	Pro	Lys	Gln	Thr	Gln	Thr	Pro	Asp	
			260					265					270			
Leu	Leu	Pro	Glu	Ala	Leu	Glu	Ala	Gln	Val	Leu	Pro	Arg	Phe	Gln	Pro	
		275					280					285				
Arg	Val	Leu	Gln	Val	Gln	Ala	Gln	Val	Gln	Ser	Gln	Thr	Gln	Pro	Arg	
	290					295					300					
Ile	Pro	Ser	Thr	Asp	Thr	Gln	Val	Gln	Pro	Lys	Leu	Gln	Lys	Gln	Ala	
305					310					315					320	
Gln	Thr	Gln	Thr	Ser	Pro	Glu	His	Leu	Val	Leu	Gln	Gln	Lys	Gln	Val	
				325					330					335		
Gln	Pro	Gln	Leu	Gln	Gln	Glu	Ala	Glu	Pro	Gln	Lys	Gln	Val	Gln	Pro	
			340					345					350			
Gln	Val	Gln	Pro	Gln	Ala	His	Ser	Gln	Gly	Pro	Arg	Gln	Val	Gln	Leu	
		355					360					365				



Gln Gln Glu Ala Glu Pro Leu Lys Gln Val Gln Pro Gln Val Gln Pro  
 370 375 380

Gln Ala His Ser Gln Pro Pro Arg Gln Val Gln Leu Gln Leu Gln Lys  
 385 390 395 400

Gln Val Gln Thr Gln Thr Tyr Pro Gln Val His Thr Gln Ala Gln Pro  
 405 410 415

Ser Val Gln Pro Gln Glu His Pro Pro Ala Gln Val Ser Val Gln Pro  
 420 425 430

Pro Glu Gln Thr His Glu Gln Pro His Thr Gln Pro Gln Val Ser Leu  
 435 440 445

Leu Ala Pro Glu Gln Thr Pro Val Val Val His Val Cys Gly Leu Glu  
 450 455 460

Met Pro Pro Asp Ala Val Glu Ala Gly Gly Gly Met Glu Lys Thr Leu  
 465 470 475 480

Pro Glu Pro Val Gly Thr Gln Val Ser Met Glu Glu Ile Gln Asn Glu  
 485 490 495

Ser Ala Cys Gly Leu Asp Val Gly Glu Cys Glu Asn Arg Ala Arg Glu  
 500 505 510

Met Pro Gly Val Trp Gly Ala Gly Gly Ser Leu Lys Val Thr Ile Leu  
 515 520 525

Gln Ser Ser Asp Ser Arg Ala Phe Ser Thr Val Pro Leu Thr Pro Val  
 530 535 540

Pro Arg Pro Ser Asp Ser Val Ser Ser Thr Pro Ala Ala Thr Ser Thr  
 545 550 555 560

Pro Ser Lys Gln Ala Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser  
 565 570 575

Cys Ser Ser Gln Gln Glu Phe Gln Asp His Met Ser Glu Pro Gln His  
 580 585 590

Gln Gln Arg Leu Gly Glu Ile Gln His Met Ser Gln Ala Cys Leu Leu

595					600					605					
Ser	Leu	Leu	Pro	Val	Pro	Arg	Asp	Val	Leu	Glu	Thr	Glu	Asp	Glu	Glu
610						615				620					
Pro	Pro	Pro	Arg	Arg	Trp	Cys	Asn	Thr	Cys	Gln	Leu	Tyr	Tyr	Met	Gly
625					630					635					640
Asp	Leu	Ile	Gln	His	Arg	Arg	Thr	Gln	Asp	His	Lys	Ile	Ala	Lys	Gln
				645					650					655	
Ser	Leu	Arg	Pro	Phe	Cys	Thr	Val	Cys	Asn	Arg	Tyr	Phe	Lys	Thr	Pro
			660					665					670		
Arg	Lys	Phe	Val	Glu	His	Val	Lys	Ser	Gln	Gly	His	Lys	Asp	Lys	Ala
		675					680					685			
Lys	Glu	Leu	Lys	Ser	Leu	Glu	Lys	Glu	Ile	Ala	Gly	Gln	Asp	Glu	Asp
	690					695					700				
His	Phe	Ile	Thr	Val	Asp	Ala	Val	Gly	Cys	Phe	Glu	Gly	Asp	Glu	Glu
705						710					715				720
Glu	Glu	Glu	Asp	Asp	Glu	Asp	Glu	Glu	Glu	Ile	Glu	Val	Glu	Glu	Glu
				725					730					735	
Leu	Cys	Lys	Gln	Val	Arg	Ser	Arg	Asp	Ile	Ser	Arg	Glu	Glu	Trp	Lys
			740					745					750		
Gly	Ser	Glu	Thr	Tyr	Ser	Pro	Asn	Thr	Ala	Tyr	Gly	Val	Asp	Phe	Leu
		755					760					765			
Val	Pro	Val	Met	Gly	Tyr	Ile	Cys	Arg	Ile	Cys	His	Lys	Phe	Tyr	His
			770				775					780			
Ser	Asn	Ser	Gly	Ala	Gln	Leu	Ser	His	Cys	Lys	Ser	Leu	Gly	His	Phe
785						790					795				800
Glu	Asn	Leu	Gln	Lys	Tyr	Lys	Ala	Ala	Lys	Asn	Pro	Ser	Pro	Thr	Thr
				805					810					815	
Arg	Pro	Val	Ser	Arg	Arg	Cys	Ala	Ile	Asn	Ala	Arg	Asn	Ala	Leu	Thr
			820					825					830		

Ala Leu Phe Thr Ser Ser Gly Arg Pro Pro Ser Gln Pro Asn Thr Gln  
835 840 845

Asp Lys Thr Pro Ser Lys Val Thr Ala Arg Pro Ser Gln Pro Pro Leu  
850 855 860

Pro Arg Arg Ser Thr Arg Leu Lys Thr  
865 870

<210> 60  
<211> 892  
<212> PRT  
<213> Homo sapiens

<400> 60

Met Phe Ser Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Leu  
1 5 10 15

Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln  
20 25 30

Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln Ala  
35 40 45

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
50 55 60

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
65 70 75 80

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Leu  
85 90 95

Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu Thr  
100 105 110

Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala Ser  
115 120 125

Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr Pro  
130 135 140

Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu Gly  
 145 150 155 160

Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser  
 165 170 175

Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro  
 180 185 190

Asn Arg Lys Thr Met Pro Val Glu Asp Lys Ser Asp Pro Pro Glu Gly  
 195 200 205

Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr Pro Glu Asp Gln Asp  
 210 215 220

Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu Lys Arg Thr Pro Ala  
 225 230 235 240

Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu Pro Ala Lys Arg Leu  
 245 250 255

Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro Pro Gly Gln Leu Gln  
 260 265 270

Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val Pro Lys Gln Thr Gln  
 275 280 285

Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala Gln Val Leu Pro Arg  
 290 295 300

Phe Gln Pro Arg Val Leu Gln Val Gln Ala Gln Val Gln Ser Gln Thr  
 305 310 315 320

Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln Val Gln Pro Lys Leu Gln  
 325 330 335

Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu His Leu Val Leu Gln Gln  
 340 345 350

Lys Gln Val Gln Pro Gln Leu Gln Gln Glu Ala Glu Pro Gln Lys Gln  
 355 360 365

Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln Gly Pro Arg Gln

370		375		380
Val Gln Leu Gln Gln Glu Ala Glu Pro Leu Lys Gln Val Gln Pro Gln				
385		390		395 400
Val Gln Pro Gln Ala His Ser Gln Pro Pro Arg Gln Val Gln Leu Gln				
	405		410	415
Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr Pro Gln Val His Thr Gln				
	420		425	430
Ala Gln Pro Ser Val Gln Pro Gln Glu His Pro Pro Ala Gln Val Ser				
	435		440	445
Val Gln Pro Pro Glu Gln Thr His Glu Gln Pro His Thr Gln Pro Gln				
	450		455	460
Val Ser Leu Leu Ala Pro Glu Gln Thr Pro Val Val Val His Val Cys				
465		470		475 480
Gly Leu Glu Met Pro Pro Asp Ala Val Glu Ala Gly Gly Gly Met Glu				
	485		490	495
Lys Thr Leu Pro Glu Pro Val Gly Thr Gln Val Ser Met Glu Glu Ile				
	500		505	510
Gln Asn Glu Ser Ala Cys Gly Leu Asp Val Gly Glu Cys Glu Asn Arg				
	515		520	525
Ala Arg Glu Met Pro Gly Val Trp Gly Ala Gly Gly Ser Leu Lys Val				
	530		535	540
Thr Ile Leu Gln Ser Ser Asp Ser Arg Ala Phe Ser Thr Val Pro Leu				
545		550		555 560
Thr Pro Val Pro Arg Pro Ser Asp Ser Val Ser Ser Thr Pro Ala Ala				
	565		570	575
Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln Phe Phe Cys Tyr Ile Cys				
	580		585	590
Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe Gln Asp His Met Ser Glu				
	595		600	605

Pro Gln His Gln Gln Arg Leu Gly Glu Ile Gln His Met Ser Gln Ala  
 610 615 620

Cys Leu Leu Ser Leu Leu Pro Val Pro Arg Asp Val Leu Glu Thr Glu  
 625 630 635 640

Asp Glu Glu Pro Pro Pro Arg Arg Trp Cys Asn Thr Cys Gln Leu Tyr  
 645 650 655

Tyr Met Gly Asp Leu Ile Gln His Arg Arg Thr Gln Asp His Lys Ile  
 660 665 670

Ala Lys Gln Ser Leu Arg Pro Phe Cys Thr Val Cys Asn Arg Tyr Phe  
 675 680 685

Lys Thr Pro Arg Lys Phe Val Glu His Val Lys Ser Gln Gly His Lys  
 690 695 700

Asp Lys Ala Lys Glu Leu Lys Ser Leu Glu Lys Glu Ile Ala Gly Gln  
 705 710 715 720

Asp Glu Asp His Phe Ile Thr Val Asp Ala Val Gly Cys Phe Glu Gly  
 725 730 735

Asp Glu Glu Glu Glu Glu Asp Asp Glu Asp Glu Glu Glu Ile Glu Val  
 740 745 750

Glu Glu Glu Leu Cys Lys Gln Val Arg Ser Arg Asp Ile Ser Arg Glu  
 755 760 765

Glu Trp Lys Gly Ser Glu Thr Tyr Ser Pro Asn Thr Ala Tyr Gly Val  
 770 775 780

Asp Phe Leu Val Pro Val Met Gly Tyr Ile Cys Arg Ile Cys His Lys  
 785 790 795 800

Phe Tyr His Ser Asn Ser Gly Ala Gln Leu Ser His Cys Lys Ser Leu  
 805 810 815

Gly His Phe Glu Asn Leu Gln Lys Tyr Lys Ala Ala Lys Asn Pro Ser  
 820 825 830

Pro Thr Thr Arg Pro Val Ser Arg Arg Cys Ala Ile Asn Ala Arg Asn  
835 840 845

Ala Leu Thr Ala Leu Phe Thr Ser Ser Gly Arg Pro Pro Ser Gln Pro  
850 855 860

Asn Thr Gln Asp Lys Thr Pro Ser Lys Val Thr Ala Arg Pro Ser Gln  
865 870 875 880

Pro Pro Leu Pro Arg Arg Ser Thr Arg Leu Lys Thr  
885 890

<210> 61  
<211> 868  
<212> PRT  
<213> Homo sapiens

<400> 61

Met Phe Ser Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Leu  
1 5 10 15

Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln  
20 25 30

Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln Ala  
35 40 45

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
50 55 60

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
65 70 75 80

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Leu  
85 90 95

Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu Thr  
100 105 110

Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala Ser  
115 120 125

Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr Pro

130		135		140
Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu Gly				
145		150		155 160
Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser				
	165		170	175
Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro				
	180		185	190
Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys Ser				
	195		200	205
Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr				
	210		215	220
Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu				
	225		230	235 240
Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu				
	245		250	255
Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro				
	260		265	270
Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val				
	275		280	285
Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala				
	290		295	300
Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala Gln				
	305		310	315 320
Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln Val				
	325		330	335
Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu His				
	340		345	350
Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu Ala				
	355		360	365



Glu Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser  
 370 375 380

Gln Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu Lys  
 385 390 395 400

Gln Val Gln Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln  
 405 410 415

Glu His Pro Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His  
 420 425 430

Glu Gln Pro His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln  
 435 440 445

Thr Pro Val Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala  
 450 455 460

Val Glu Ala Gly Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly  
 465 470 475 480

Thr Gln Val Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu  
 485 490 495

Asp Val Gly Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp  
 500 505 510

Gly Ala Gly Gly Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser  
 515 520 525

Arg Ala Phe Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp  
 530 535 540

Ser Val Ser Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala  
 545 550 555 560

Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln  
 565 570 575

Glu Phe Gln Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly  
 580 585 590

Glu Ile Gln His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro Val  
 595 600 605

Pro Arg Asp Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg  
 610 615 620

Trp Cys Asn Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His  
 625 630 635 640

Arg Arg Thr Gln Asp His Lys Ile Ala Lys Gln Ser Leu Arg Pro Phe  
 645 650 655

Cys Thr Val Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu  
 660 665 670

His Val Lys Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser  
 675 680 685

Leu Glu Lys Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val  
 690 695 700

Asp Ala Val Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Asp Asp  
 705 710 715 720

Glu Asp Glu Glu Glu Ile Glu Val Glu Glu Glu Leu Cys Lys Gln Val  
 725 730 735

Arg Ser Arg Asp Ile Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr Tyr  
 740 745 750

Ser Pro Asn Thr Ala Tyr Gly Val Asp Phe Leu Val Pro Val Met Gly  
 755 760 765

Tyr Ile Cys Arg Ile Cys His Lys Phe Tyr His Ser Asn Ser Gly Ala  
 770 775 780

Gln Leu Ser His Cys Lys Ser Leu Gly His Phe Glu Asn Leu Gln Lys  
 785 790 795 800

Tyr Lys Ala Ala Lys Asn Pro Ser Pro Thr Thr Arg Pro Val Ser Arg  
 805 810 815

Arg Cys Ala Ile Asn Ala Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser  
820 825 830

Ser Gly Arg Pro Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro Ser  
835 840 845

Lys Val Thr Ala Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser Thr  
850 855 860

Arg Leu Lys Thr  
865

<210> 62  
<211> 841  
<212> PRT  
<213> Homo sapiens

<400> 62

Met Phe Ser Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Leu  
1 5 10 15

Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln  
20 25 30

Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln Ala  
35 40 45

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
50 55 60

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
65 70 75 80

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Leu  
85 90 95

Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu Thr  
100 105 110

Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala Ser  
115 120 125

Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr Pro  
130 135 140

Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu Gly  
 145 150 155 160

Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser  
 165 170 175

Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro  
 180 185 190

Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys Ser  
 195 200 205

Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr  
 210 215 220

Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu  
 225 230 235 240

Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu  
 245 250 255

Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro  
 260 265 270

Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val  
 275 280 285

Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala  
 290 295 300

Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala Gln  
 305 310 315 320

Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln Val  
 325 330 335

Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu His  
 340 345 350

Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu Ala  
 355 360 365

Glu Pro Gln Lys Gln Val Gln Pro Gln Val His Thr Gln Ala Gln Pro  
 370 375 380

Ser Val Gln Pro Gln Glu His Pro Pro Ala Gln Val Ser Val Gln Pro  
 385 390 395 400

Pro Glu Gln Thr His Glu Gln Pro His Thr Gln Pro Gln Val Ser Leu  
 405 410 415

Leu Ala Pro Glu Gln Thr Pro Val Val Val His Val Cys Gly Leu Glu  
 420 425 430

Met Pro Pro Asp Ala Val Glu Ala Gly Gly Gly Met Glu Lys Thr Leu  
 435 440 445

Pro Glu Pro Val Gly Thr Gln Val Ser Met Glu Glu Ile Gln Asn Glu  
 450 455 460

Ser Ala Cys Gly Leu Asp Val Gly Glu Cys Glu Asn Arg Ala Arg Glu  
 465 470 475 480

Met Pro Gly Val Trp Gly Ala Gly Gly Ser Leu Lys Val Thr Ile Leu  
 485 490 495

Gln Ser Ser Asp Ser Arg Ala Phe Ser Thr Val Pro Leu Thr Pro Val  
 500 505 510

Pro Arg Pro Ser Asp Ser Val Ser Ser Thr Pro Ala Ala Thr Ser Thr  
 515 520 525

Pro Ser Lys Gln Ala Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser  
 530 535 540

Cys Ser Ser Gln Gln Glu Phe Gln Asp His Met Ser Glu Pro Gln His  
 545 550 555 560

Gln Gln Arg Leu Gly Glu Ile Gln His Met Ser Gln Ala Cys Leu Leu  
 565 570 575

Ser Leu Leu Pro Val Pro Arg Asp Val Leu Glu Thr Glu Asp Glu Glu  
 580 585 590

Pro Pro Pro Arg Arg Trp Cys Asn Thr Cys Gln Leu Tyr Tyr Met Gly  
 595 600 605

Asp Leu Ile Gln His Arg Arg Thr Gln Asp His Lys Ile Ala Lys Gln  
 610 615 620

Ser Leu Arg Pro Phe Cys Thr Val Cys Asn Arg Tyr Phe Lys Thr Pro  
 625 630 635 640

Arg Lys Phe Val Glu His Val Lys Ser Gln Gly His Lys Asp Lys Ala  
 645 650 655

Lys Glu Leu Lys Ser Leu Glu Lys Glu Ile Ala Gly Gln Asp Glu Asp  
 660 665 670

His Phe Ile Thr Val Asp Ala Val Gly Cys Phe Glu Gly Asp Glu Glu  
 675 680 685

Glu Glu Glu Asp Asp Glu Asp Glu Glu Glu Ile Glu Val Glu Glu Glu  
 690 695 700

Leu Cys Lys Gln Val Arg Ser Arg Asp Ile Ser Arg Glu Glu Trp Lys  
 705 710 715 720

Gly Ser Glu Thr Tyr Ser Pro Asn Thr Ala Tyr Gly Val Asp Phe Leu  
 725 730 735

Val Pro Val Met Gly Tyr Ile Cys Arg Ile Cys His Lys Phe Tyr His  
 740 745 750

Ser Asn Ser Gly Ala Gln Leu Ser His Cys Lys Ser Leu Gly His Phe  
 755 760 765

Glu Asn Leu Gln Lys Tyr Lys Ala Ala Lys Asn Pro Ser Pro Thr Thr  
 770 775 780

Arg Pro Val Ser Arg Arg Cys Ala Ile Asn Ala Arg Asn Ala Leu Thr  
 785 790 795 800

Ala Leu Phe Thr Ser Ser Gly Arg Pro Pro Ser Gln Pro Asn Thr Gln  
 805 810 815

Asp Lys Thr Pro Ser Lys Val Thr Ala Arg Pro Ser Gln Pro Pro Leu

820

825

830

Pro Arg Arg Ser Thr Arg Leu Lys Thr  
           835                  840

&lt;210&gt; 63

&lt;211&gt; 785

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 63

Met Phe Ser Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln Leu  
   1                  5                  10                  15

Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln  
           20                  25                  30

Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln Ala  
           35                  40                  45

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
   50                  55                  60

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
   65                  70                  75                  80

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Leu  
           85                  90                  95

Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu Thr  
           100                  105                  110

Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala Ser  
           115                  120                  125

Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr Pro  
           130                  135                  140

Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu Gly  
   145                  150                  155                  160

Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser  
           165                  170                  175

Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro  
 180 185 190

Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys Ser  
 195 200 205

Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr  
 210 215 220

Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu  
 225 230 235 240

Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu  
 245 250 255

Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro  
 260 265 270

Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val  
 275 280 285

Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala  
 290 295 300

Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala Pro  
 305 310 315 320

Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln Glu His Pro  
 325 330 335

Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His Glu Gln Pro  
 340 345 350

His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln Thr Pro Val  
 355 360 365

Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala Val Glu Ala  
 370 375 380

Gly Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly Thr Gln Val  
 385 390 395 400



Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu Asp Val Gly  
 405 410 415

Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp Gly Ala Gly  
 420 425 430

Gly Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser Arg Ala Phe  
 435 440 445

Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp Ser Val Ser  
 450 455 460

Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln Phe  
 465 470 475 480

Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe Gln  
 485 490 495

Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly Glu Ile Gln  
 500 505 510

His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro Val Pro Arg Asp  
 515 520 525

Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg Trp Cys Asn  
 530 535 540

Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His Arg Arg Thr  
 545 550 555 560

Gln Asp His Lys Ile Ala Lys Gln Ser Leu Arg Pro Phe Cys Thr Val  
 565 570 575

Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu His Val Lys  
 580 585 590

Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser Leu Glu Lys  
 595 600 605

Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val Asp Ala Val  
 610 615 620

Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp Asp Glu Asp Glu

625		630		635		640
Glu Glu Ile Glu Val Glu Glu Glu Leu Cys Lys Gln Val Arg Ser Arg						
	645		650		655	
Asp Ile Ser Arg Glu Glu Trp Lys Gly Ser Glu Thr Tyr Ser Pro Asn						
	660		665		670	
Thr Ala Tyr Gly Val Asp Phe Leu Val Pro Val Met Gly Tyr Ile Cys						
	675		680		685	
Arg Ile Cys His Lys Phe Tyr His Ser Asn Ser Gly Ala Gln Leu Ser						
	690		695		700	
His Cys Lys Ser Leu Gly His Phe Glu Asn Leu Gln Lys Tyr Lys Ala						
	705		710		715	
Ala Lys Asn Pro Ser Pro Thr Thr Arg Pro Val Ser Arg Arg Cys Ala						
	725		730		735	
Ile Asn Ala Arg Asn Ala Leu Thr Ala Leu Phe Thr Ser Ser Gly Arg						
	740		745		750	
Pro Pro Ser Gln Pro Asn Thr Gln Asp Lys Thr Pro Ser Lys Val Thr						
	755		760		765	
Ala Arg Pro Ser Gln Pro Pro Leu Pro Arg Arg Ser Thr Arg Leu Lys						
	770		775		780	

Thr  
785

<210> 64  
 <211> 889  
 <212> PRT  
 <213> Homo sapiens

<400> 64

Met Phe Ser Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln Leu
1 5 10 15

Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln
20 25 30

Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln Ala  
 35 40 45

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
 50 55 60

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
 65 70 75 80

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Leu  
 85 90 95

Asp Gln Phe Ala Met Pro Pro Ala Thr Tyr Asp Thr Ala Gly Leu Thr  
 100 105 110

Met Pro Thr Ala Thr Leu Gly Asn Leu Arg Gly Tyr Gly Met Ala Ser  
 115 120 125

Pro Gly Leu Ala Ala Pro Ser Leu Thr Pro Pro Gln Leu Ala Thr Pro  
 130 135 140

Asn Leu Gln Gln Phe Phe Pro Gln Ala Thr Arg Gln Ser Leu Leu Gly  
 145 150 155 160

Pro Pro Pro Val Gly Val Pro Met Asn Pro Ser Gln Phe Asn Leu Ser  
 165 170 175

Gly Arg Asn Pro Gln Lys Gln Ala Arg Thr Ser Ser Ser Thr Thr Pro  
 180 185 190

Asn Arg Lys Asp Ser Ser Ser Gln Thr Met Pro Val Glu Asp Lys Ser  
 195 200 205

Asp Pro Pro Glu Gly Ser Glu Glu Ala Ala Glu Pro Arg Met Asp Thr  
 210 215 220

Pro Glu Asp Gln Asp Leu Pro Pro Cys Pro Glu Asp Ile Ala Lys Glu  
 225 230 235 240

Lys Arg Thr Pro Ala Pro Glu Pro Glu Pro Cys Glu Ala Ser Glu Leu  
 245 250 255

Pro Ala Lys Arg Leu Arg Ser Ser Glu Glu Pro Thr Glu Lys Glu Pro  
 260 265 270

Pro Gly Gln Leu Gln Val Lys Ala Gln Pro Gln Ala Arg Met Thr Val  
 275 280 285

Pro Lys Gln Thr Gln Thr Pro Asp Leu Leu Pro Glu Ala Leu Glu Ala  
 290 295 300

Gln Val Leu Pro Arg Phe Gln Pro Arg Val Leu Gln Val Gln Ala Gln  
 305 310 315 320

Val Gln Ser Gln Thr Gln Pro Arg Ile Pro Ser Thr Asp Thr Gln Val  
 325 330 335

Gln Pro Lys Leu Gln Lys Gln Ala Gln Thr Gln Thr Ser Pro Glu His  
 340 345 350

Leu Val Leu Gln Gln Lys Gln Val Gln Pro Gln Leu Gln Gln Glu Ala  
 355 360 365

Glu Pro Gln Lys Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser  
 370 375 380

Gln Gly Pro Arg Gln Val Gln Leu Gln Gln Glu Ala Glu Pro Leu Lys  
 385 390 395 400

Gln Val Gln Pro Gln Val Gln Pro Gln Ala His Ser Gln Pro Pro Arg  
 405 410 415

Gln Val Gln Leu Gln Leu Gln Lys Gln Val Gln Thr Gln Thr Tyr Pro  
 420 425 430

Gln Val His Thr Gln Ala Gln Pro Ser Val Gln Pro Gln Glu His Pro  
 435 440 445

Pro Ala Gln Val Ser Val Gln Pro Pro Glu Gln Thr His Glu Gln Pro  
 450 455 460

His Thr Gln Pro Gln Val Ser Leu Leu Ala Pro Glu Gln Thr Pro Val  
 465 470 475 480

Val Val His Val Cys Gly Leu Glu Met Pro Pro Asp Ala Val Glu Ala

485	490	495
Gly Gly Gly Met Glu Lys Thr Leu Pro Glu Pro Val Gly Thr Gln Val 500 505 510		
Ser Met Glu Glu Ile Gln Asn Glu Ser Ala Cys Gly Leu Asp Val Gly 515 520 525		
Glu Cys Glu Asn Arg Ala Arg Glu Met Pro Gly Val Trp Gly Ala Gly 530 535 540		
Gly Ser Leu Lys Val Thr Ile Leu Gln Ser Ser Asp Ser Arg Ala Phe 545 550 555 560		
Ser Thr Val Pro Leu Thr Pro Val Pro Arg Pro Ser Asp Ser Val Ser 565 570 575		
Ser Thr Pro Ala Ala Thr Ser Thr Pro Ser Lys Gln Ala Leu Gln Phe 580 585 590		
Phe Cys Tyr Ile Cys Lys Ala Ser Cys Ser Ser Gln Gln Glu Phe Gln 595 600 605		
Asp His Met Ser Glu Pro Gln His Gln Gln Arg Leu Gly Glu Ile Gln 610 615 620		
His Met Ser Gln Ala Cys Leu Leu Ser Leu Leu Pro Val Pro Arg Asp 625 630 635 640		
Val Leu Glu Thr Glu Asp Glu Glu Pro Pro Pro Arg Arg Trp Cys Asn 645 650 655		
Thr Cys Gln Leu Tyr Tyr Met Gly Asp Leu Ile Gln His Arg Arg Thr 660 665 670		
Gln Asp His Lys Ile Ala Lys Gln Ser Leu Arg Pro Phe Cys Thr Val 675 680 685		
Cys Asn Arg Tyr Phe Lys Thr Pro Arg Lys Phe Val Glu His Val Lys 690 695 700		
Ser Gln Gly His Lys Asp Lys Ala Lys Glu Leu Lys Ser Leu Glu Lys 705 710 715 720		

Glu Ile Ala Gly Gln Asp Glu Asp His Phe Ile Thr Val Asp Ala Val  
725 730 735

Gly Cys Phe Glu Gly Asp Glu Glu Glu Glu Glu Asp Asp Glu Asp Glu  
740 745 750

Glu Glu Ile Glu Val Arg Ser Arg Asp Ile Ser Arg Glu Glu Trp Lys  
755 760 765

Gly Ser Glu Thr Tyr Ser Pro Asn Thr Ala Tyr Gly Val Asp Phe Leu  
770 775 780

Val Pro Val Met Gly Tyr Ile Cys Arg Ile Cys His Lys Phe Tyr His  
785 790 795 800

Ser Asn Ser Gly Ala Gln Leu Ser His Cys Lys Ser Leu Gly His Phe  
805 810 815

Glu Asn Leu Gln Lys Tyr Lys Ala Ala Lys Asn Pro Ser Pro Thr Thr  
820 825 830

Arg Pro Val Ser Arg Arg Cys Ala Ile Asn Ala Arg Asn Ala Leu Thr  
835 840 845

Ala Leu Phe Thr Ser Ser Gly Arg Pro Pro Ser Gln Pro Asn Thr Gln  
850 855 860

Asp Lys Thr Pro Ser Lys Val Thr Ala Arg Pro Ser Gln Pro Pro Leu  
865 870 875 880

Pro Arg Arg Ser Thr Arg Leu Lys Thr  
885

<210> 65  
<211> 873  
<212> PRT  
<213> Homo sapiens

<400> 65

Met Phe Ser Gln Gln Gln Gln Gln Leu Gln Gln Gln Gln Gln Gln Leu  
1 5 10 15

Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln Gln Leu Gln Gln Gln  
 20 25 30

Gln Leu Leu Gln Leu Gln Gln Leu Leu Gln Gln Ser Pro Pro Gln Ala  
 35 40 45

Pro Leu Pro Met Ala Val Ser Arg Gly Leu Pro Pro Gln Gln Pro Gln  
 50 55 60

Gln Pro Leu Leu Asn Leu Gln Gly Thr Asn Ser Ala Ser Leu Leu Asn  
 65 70 75 80

Gly Ser Met Leu Gln Arg Ala Leu Leu Leu Gln Gln Leu Gln Gly Asn  
 85 90 95

Leu Arg Gly Tyr Gly Met Ala Ser Pro Gly Leu Ala Ala Pro Ser Leu  
 100 105 110

Thr Pro Pro Gln Leu Ala Thr Pro Asn Leu Gln Gln Phe Phe Pro Gln  
 115 120 125

Ala Thr Arg Gln Ser Leu Leu Gly Pro Pro Pro Val Gly Val Pro Met  
 130 135 140

Asn Pro Ser Gln Phe Asn Leu Ser Gly Arg Asn Pro Gln Lys Gln Ala  
 145 150 155 160

Arg Thr Ser Ser Ser Thr Thr Pro Asn Arg Lys Asp Ser Ser Ser Gln  
 165 170 175

Thr Met Pro Val Glu Asp Lys Ser Asp Pro Pro Glu Gly Ser Glu Glu  
 180 185 190

Ala Ala Glu Pro Arg Met Asp Thr Pro Glu Asp Gln Asp Leu Pro Pro  
 195 200 205

Cys Pro Glu Asp Ile Ala Lys Glu Lys Arg Thr Pro Ala Pro Glu Pro  
 210 215 220

Glu Pro Cys Glu Ala Ser Glu Leu Pro Ala Lys Arg Leu Arg Ser Ser  
 225 230 235 240

Glu Glu Pro Thr Glu Lys Glu Pro Pro Gly Gln Leu Gln Val Lys Ala





Pro Glu Pro Val Gly Thr Gln Val Ser Met Glu Glu Ile Gln Asn Glu  
 485 490 495

Ser Ala Cys Gly Leu Asp Val Gly Glu Cys Glu Asn Arg Ala Arg Glu  
 500 505 510

Met Pro Gly Val Trp Gly Ala Gly Gly Ser Leu Lys Val Thr Ile Leu  
 515 520 525

Gln Ser Ser Asp Ser Arg Ala Phe Ser Thr Val Pro Leu Thr Pro Val  
 530 535 540

Pro Arg Pro Ser Asp Ser Val Ser Ser Thr Pro Ala Ala Thr Ser Thr  
 545 550 555 560

Pro Ser Lys Gln Ala Leu Gln Phe Phe Cys Tyr Ile Cys Lys Ala Ser  
 565 570 575

Cys Ser Ser Gln Gln Glu Phe Gln Asp His Met Ser Glu Pro Gln His  
 580 585 590

Gln Gln Arg Leu Gly Glu Ile Gln His Met Ser Gln Ala Cys Leu Leu  
 595 600 605

Ser Leu Leu Pro Val Pro Arg Asp Val Leu Glu Thr Glu Asp Glu Glu  
 610 615 620

Pro Pro Pro Arg Arg Trp Cys Asn Thr Cys Gln Leu Tyr Tyr Met Gly  
 625 630 635 640

Asp Leu Ile Gln His Arg Arg Thr Gln Asp His Lys Ile Ala Lys Gln  
 645 650 655

Ser Leu Arg Pro Phe Cys Thr Val Cys Asn Arg Tyr Phe Lys Thr Pro  
 660 665 670

Arg Lys Phe Val Glu His Val Lys Ser Gln Gly His Lys Asp Lys Ala  
 675 680 685

Lys Glu Leu Lys Ser Leu Glu Lys Glu Ile Ala Gly Gln Asp Glu Asp  
 690 695 700

His Phe Ile Thr Val Asp Ala Val Gly Cys Phe Glu Gly Asp Glu Glu  
705 710 715 720

Glu Glu Glu Asp Asp Glu Asp Glu Glu Glu Ile Glu Val Glu Glu Glu  
725 730 735

Leu Cys Lys Gln Val Arg Ser Arg Asp Ile Ser Arg Glu Glu Trp Lys  
740 745 750

Gly Ser Glu Thr Tyr Ser Pro Asn Thr Ala Tyr Gly Val Asp Phe Leu  
755 760 765

Val Pro Val Met Gly Tyr Ile Cys Arg Ile Cys His Lys Phe Tyr His  
770 775 780

Ser Asn Ser Gly Ala Gln Leu Ser His Cys Lys Ser Leu Gly His Phe  
785 790 795 800

Glu Asn Leu Gln Lys Tyr Lys Ala Ala Lys Asn Pro Ser Pro Thr Thr  
805 810 815

Arg Pro Val Ser Arg Arg Cys Ala Ile Asn Ala Arg Asn Ala Leu Thr  
820 825 830

Ala Leu Phe Thr Ser Ser Gly Arg Pro Pro Ser Gln Pro Asn Thr Gln  
835 840 845

Asp Lys Thr Pro Ser Lys Val Thr Ala Arg Pro Ser Gln Pro Pro Leu  
850 855 860

Pro Arg Arg Ser Thr Arg Leu Lys Thr  
865 870

<210> 66  
<211> 2821  
<212> DNA  
<213> Homo sapiens

<400> 66  
tgggggctgc ggggccggcc catccgtggg ggcgacttga gcgttgaggg cgcgcgggga 60  
ggcgagccac catgttcagc cagcagcagc agcagctcca gcaacagcag ggccccgttg 120  
cccatggctg tcagccgggg gctccccccg cagcagccac agcagccgct tctgaatctc 180

cagggcacca	actcagcctc	cctcctcaac	ggctccatgc	tgcagagagc	tttgctttta	240
cagcagttgc	aaggactgga	ccagtttgca	atgccaccag	ccacgtatga	cactgccggg	300
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ctcgcagccc	ccagcctcac	acccccacaa	ctggccactc	caaatttgca	acagttcttt	420
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